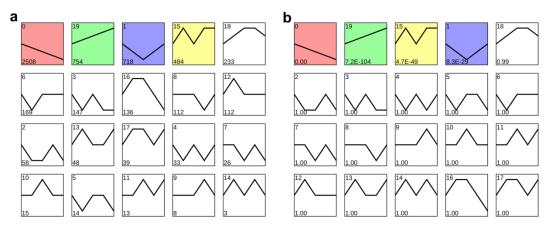
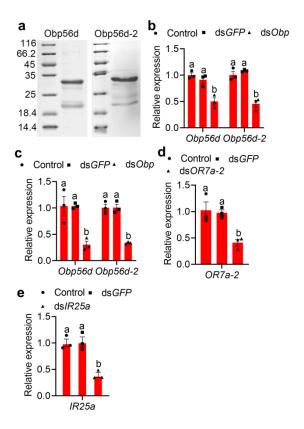


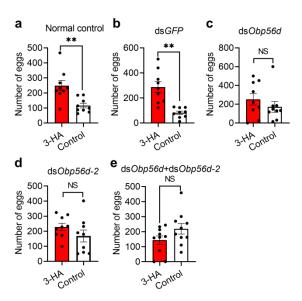
Supplementary Figure 1 Comparison of number of eggs laid in purees and purees mixed with 3-HA. (a) Eggs laid in guava purees and purees mixed with 3-HA (n = 13 replicates, Kendall W = 0.28,  $\chi^2 = 10.938$ , P = 0.012). (b) Eggs laid in mango purees and purees mixed with 3-HA (n = 23 replicates, Kendall W = 0.136,  $\chi^2 = 9.393$ , P = 0.024). Number of blue dots in each bar indicated the replicates. Different letters above the error bars indicate significant differences at the 0.05 level with the Kendall nonparametric test. Data in bar plots show mean values  $\pm$  SEM.



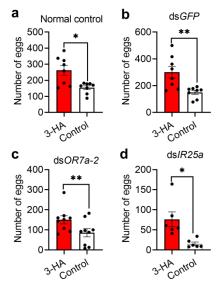
Supplementary Figure 2 Number of genes in different expression patterns (a) and statistical analysis (b).



Supplementary Figure 3 Gene expressions analysis. (a) Recombinant protein of Obp56d and Obp56d-2 verification. (b) Expression of Obp56d and Obp56d-2 gene 24 h after their dsRNA was injected individually (Obp56d: F = 20.07, df = 2, 6, P = 0.0022; Obp56d-2: F = 39.16, df = 2, 6, P = 0.0004). (c) Expression of Obp56d and Obp56d-2 gene 24 h after their dsRNA was injected together (Obp56d: F = 13.61, df = 2, 6, P = 0.0059; Obp56d-2: F = 51.5, df = 2, 6, P = 0.0002). (d) Expression of OR7a-2 gene 24 h after its dsRNA was injected (F = 10.78, df = 2, 6, P = 0.0103). (e) Expression of IR25a gene 24 h after its dsRNA was injected (F = 23.83, df = 2, 6, P = 0.0014). Number of black dots, squares and triangles in each bar indicated the replicates (F = 3 replicates). Different letters above the error bars indicate significant differences at the 0.05 level analyzed by ANOVA followed by Tukey's test. Data in bar plots show mean values  $\pm$  SEM. Detail information for sample sizes can be found in Supplementary Data 2.



Supplementary Figure 4 Comparison of egg number in guava puree and guava puree added 3-HA. (a) Comparison of the number of eggs laid by normal flies in guava puree and guava puree added 3-HA (n = 9 replicates, P = 0.0039, Sum of signed ranks (W) = -45). (b) Comparison of the number of eggs laid by dsGFP injected flies in guava puree and guava puree added 3-HA (n = 9 replicates, P = 0.0039, Sum of signed ranks (W) = -45). (c) Comparison of the number of eggs laid by dsObp56d injected flies in guava puree and guava puree added 3-HA (n = 9 replicates, P = 0.375, Sum of signed ranks (W) = -16). (d) Comparison of the number of eggs laid by dsObp56d-2 injected flies in guava puree and guava puree added 3-HA (n = 9 replicates, P = 0.25, Sum of signed ranks (W) = -21). (e) Comparison of the number of eggs laid by dsObp56d-and dsObp56d-2 injected flies in guava puree and guava puree added 3-HA (n = 10 replicates, p = 0.4922, Sum of signed ranks (W) = 15). The data was analyzed with Wilcoxon matched-pairs signed rank test. "NS" no significance, \*\* P < 0.01. Data in bar plots show mean values  $\pm$  SEM.



Supplementary Figure 5 Comparison of egg number in guava puree and guava puree added 3-HA. (a) Comparison of the number of eggs laid by normal flies in

guava puree and guava puree added 3-HA (n=8 replicates, P=0.0156, Sum of signed ranks (W) = -34). (b) Comparison of the number of eggs laid by dsGFP injected flies in guava puree and guava puree added 3-HA (n=8 replicates, P=0.0078, Sum of signed ranks (W) = -36). (c) Comparison of the number of eggs laid by dsOR7a-2 injected flies in guava puree and guava puree added 3-HA (n=9 replicates, P=0.0078, Sum of signed ranks (W) = -36). (d) Comparison of the number of eggs laid by dsIR25a injected flies in guava puree and guava puree added 3-HA (n=6 replicates, P=0.0313, Sum of signed ranks (W) = -21). The data was analyzed with Wilcoxon matched-pairs signed rank test. \* P<0.05, \*\* P<0.01. Data in bar plots show mean values  $\pm$  SEM.

Supplementary Table 1 Primers used in qRT-PCR experimental

| Primer name         | Sequence (5'-3')     | Length (bp) |
|---------------------|----------------------|-------------|
| α-TUB-F             | CGCATTCATGGTTGATAACG | 184         |
| $\alpha$ - $TUB$ -R | GGGCACCAAGTTAGTCTGGA |             |
| ACT5-F              | CAACTCACCCGCAATGTATG | 237         |
| ACT5-R              | CGCTCAGCAGTGGTTGTAAA |             |
| Obp19d-F            | TAGTGGTTTTACCTGCGCGT | 92          |
| Obp19d-R            | AGCAAAGCTGCAGATAGCCA |             |
| <i>Obp56d-2-</i> F  | TTCCGAAGAGGCGATTCGTG | 88          |
| <i>Obp56d-2-</i> R  | CTCTTGCATGCACTTGGAGA |             |
| Obp56d-F            | TTCGCTGTTGCTGTTTTGCT | 99          |
| Obp56d-R            | TGCAGCGTGTACCTTTTGTT |             |
| <i>OR7a-2-</i> F    | CTTTCTGTGCTACTCGGCCA | 112         |
| <i>OR7a-2-</i> R    | TATAAACTCCGTGCGGGAGC |             |
| <i>IR25a</i> -F     | ATGCAGGAAGCCCAGTTACC | 71          |
| <i>IR25a</i> -R     | GTGGCTGAAGTTGAGTTGCG |             |

## Supplementary Table 2 Primers used for recombinant proteins and dsRNA synthesis

| Sequence (5'-3')       |   |  |
|------------------------|---|--|
| Re-Obp56d-F            | GGTGGTGGATCCGAATTCCGGACTATGCAGGAAGGCGTTGGCAAACTGACCGAAGAACA |  |
| Re-Obp56d-R            | GGTGGTGCTCGAGTGCGGCCTTATTAAAACACAAAGCTTTCATGGCCTGCTGCATA    |  |
| Re- <i>Obp56d</i> -2 F | GGTGGTGGATCCGAATTCCGGACTATGCAGGAAGGCGTTAATAAACTGACCGAAGAACA |  |
| Re- <i>Obp56d</i> -2 R | GGTGGTGCTCGAGTGCGGCCTTATTAAAAATCCAGGCCTTCATGACCGGCGCAT      |  |
| ds <i>Obp19d-</i> F    | TAATACGACTCACTATAGTTCGCTGTTGCTGTTTTGCTA                     |  |
| ds <i>Obp19d-</i> R    | TAATACGACTCACTATAGTAAGGCCTTTGGACTTCGCA                      |  |
| ds <i>Obp56d-</i> 2 F  | TAATACGACTCACTATAGTTCGCTGTTGCTGTTTTGCTA                     |  |
| ds <i>Obp56d-</i> 2 R  | TAATACGACTCACTATAGATCCAAGCCCTCGTGCC                         |  |
| ds <i>Obp56d-</i> F    | TAATACGACTCACTATAGTTCGCTGTTGCTGTTTTGCTA                     |  |
| ds <i>Obp56d-</i> R    | TAATACGACTCACTATAGAAAACAAAGCTCTCATGCCCA                     |  |
| ds <i>OR7a-2-</i> F    | TAATACGACTCACTATAGCTACCACTTGCCGATTGATGC                     |  |
| ds <i>OR7a-2-</i> R    | TAATACGACTCACTATAGCGAAGGCATGACTAAGCCCA                      |  |
| ds <i>IR25a-</i> F     | TAATACGACTCACTATAGAGCCCCTACAGCTACCAGAA                      |  |
| ds <i>IR25a</i> -R     | TAATACGACTCACTATAGCACCGGTGTATCGAGACGAG                      |  |

Note: The blue sequence is the homologous sequence of vector; The red sequence is T7 promoter.